Documentation of `ModelSolver` [*]
A `Python` class for analyzing dynamic algebraic models

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Abstract

This paper documents the `Python` class `ModelSolver`. `ModelSolver` lets the user define a model object in terms of equations and endogenous variables. It contains methods to solve the model subject to data (in a `Pandas DataFrame`), as well as analyzing the model using graph theory and network plots.

What sets `ModelSolver` apart from other similar solvers, is that it does not require the equations of the model to be written in any particular form, or that the user associates equations with endogenous variables. Most other solvers require that either 1) the model is normalized (i.e., that the model is written in terms of endogenous variables), 2) or that the user explicitly associates equations with endogenous variables. This is non-trivial for models with lots of equations. `ModelSolver`, however, reads equations in whatever form they may be formulated, and performs the necessary analyzes, without any input from the user other than lists of equations and endogenous variables.

`ModelSolver` was developed in order to facilitate solving a monthly input-output model for the Norwegian monthly national accounts. It analyzes and solves the model's more than 15,500 equations for more than 30 periods in under a minute on a laptop.

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[*]https://github.com/statisticsnorway/model-solver.git
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1 Background

1.1 The model

Suppose we have a model consisting of n equations and endogenous variables,
[1]

$$L_1(\mathbf{x}_t, \mathbf{z}_t) = H_1(\mathbf{x}_t, \mathbf{z}_t),$$
$$L_2(\mathbf{x}_t, \mathbf{z}_t) = H_2(\mathbf{x}_t, \mathbf{z}_t),$$
$$\vdots$$
$$L_n(\mathbf{x}_t, \mathbf{z}_t) = H_n(\mathbf{x}_t, \mathbf{z}_t),$$

for $t = T_0, T_0 + 1, \ldots, T_1$, where $\mathbf{x}_t = (x_{1,t}, x_{2,t}, \ldots, x_{n,t})$ is a vector of endogenous variables, and $\mathbf{z}_t = (z_{1,t-k_1}, z_{2,t-k_2}, \ldots, z_{m,t-k_m}, \ldots, x_{i,t-\kappa_i}, \ldots)$, for $k_i \geq 0$ and $\kappa_i > 0$, is a vector of simultaneous and lags of exogenous variables and lags of endogenous variables. [2] It's convenient to formulate the model on vector form,

$$\mathbf{F}(\mathbf{x}_t, \mathbf{z}_t) = \begin{pmatrix} F_1(\mathbf{x}_t, \mathbf{z}_t) \\ F_2(\mathbf{x}_t, \mathbf{z}_t) \\ \vdots \\ F_n(\mathbf{x}_t, \mathbf{z}_t) \end{pmatrix} = \begin{pmatrix} L_1(\mathbf{x}_t, \mathbf{z}_t) - H_1(\mathbf{x}_t, \mathbf{z}_t) \\ L_2(\mathbf{x}_t, \mathbf{z}_t) - H_2(\mathbf{x}_t, \mathbf{z}_t) \\ \vdots \\ L_n(\mathbf{x}_t, \mathbf{z}_t) - H_n(\mathbf{x}_t, \mathbf{z}_t) \end{pmatrix},$$

with the solution to the model being given by $\{\mathbf{x}_t\}_{t=T_0}^{T_1}$ such that

$$\mathbf{F}(\mathbf{x}_t, \mathbf{z}_t) = \mathbf{0} \text{ for } t = T_0, T_0 + 1, \ldots, T_1.$$

On the face of it, this is a problem which the *Newton-Raphson*-algorithm handles well. The issue, however, is that n might be quite large. In the Norwegian national accounts, for instance, n is just greater than 15,500. Therefore, it is useful to analyze the system of equations before solving it, in order to break it down into minimal *simultaneous blocks* that can be solved in a particular sequence.

1.2 Block analysis

In order to analyze and divide the model into blocks, we use results from *graph theory*.

First, we construct a *bipartite graph* (BiGraph) that connects endogenous variables with equations. This is illustrated in Figure 1 for an arbitrary model with 4 equations (we omit time subscripts and exogenous variables and lags for notational convenience).

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[1] L_i and H_i denote the left and right hand side of equation i.
[2] Note that $\mathbf{z}_t$ may contain only $z_{i,t-k_i}$'s or $x_{i,t-k_i}$'s or be empty.
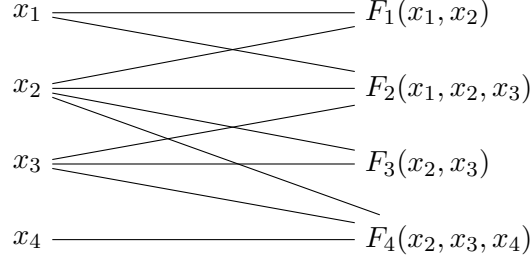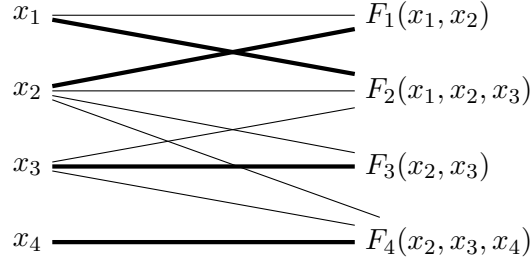
Figure 1: Bipartite graph (BiGraph of model)

$$x_1 \quad\text{---}\quad F_1(x_1, x_2)$$
$$x_2 \quad\text{---}\quad F_2(x_1, x_2, x_3)$$
$$x_3 \quad\text{---}\quad F_3(x_2, x_3)$$
$$x_4 \quad\text{---}\quad F_4(x_2, x_3, x_4)$$

Figure 2: Maximum Bipartite Match (MBM) of BiGraph

$$x_1 \quad\text{---}\quad F_1(x_1, x_2)$$
$$x_2 \quad\text{---}\quad F_2(x_1, x_2, x_3)$$
$$x_3 \quad\text{---}\quad F_3(x_2, x_3)$$
$$x_4 \quad\text{---}\quad F_4(x_2, x_3, x_4)$$

Next, we apply *maximum bipartite matching* (MBM), which assigns each endogenous variable to *one and only one* equation. The MBM is arbitrary, and Figure 2 shows one possible solution.

We use Figure 2 to determine what endogenous variables impact what *other* endogenous variables. This is illustrated in Figure 3.

Next, we use Figure 3 to construct a *directed graph* (DiGraph) that shows how the endogenous variables impact each other. The DiGraph is shown in Figure 4.

Finally, we find the *strong components* of the DiGraph, as shown in Figure 5. A strong component is a set of nodes that are connected such that every node can be reached from every other node in the set (traversing the arrows). A *condensation* of the DiGraph is a (new) DiGraph with each node being a strong component (of the former). The condensation can be illustrated as in Figure 6. Each node of the condensation corresponds to a block of the model, and the arrows decide the sequence in which the blocks are to be solved. See [1] for a discussion of strong components and block analysis.

In this example, x_1 and x_2 must be solved first in one simultaneous block. Next, x_3 is solved (taking x_1 and x_2 as given, being predetermined by the solution to block 1). Finally, x_4 is solved (taking x_1, x_2 and x_3 as given, being predetermined by the solutions to block 1 and 2).

Figure 3: Graph of what endogenous variables impact what *other* endogenous variables
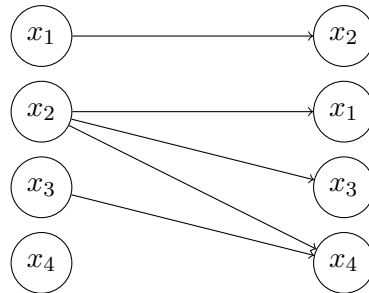

Figure 4: Directed graph (DiGraph) of what endogenous variables impact what *other* endogenous variables
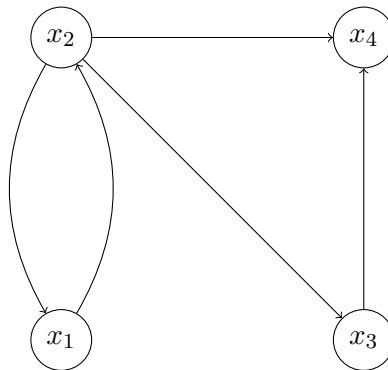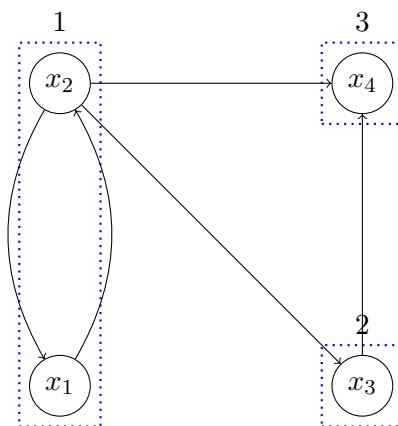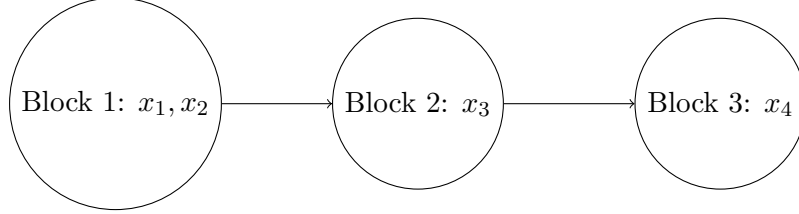

Figure 5: Condensation of DiGraph

Figure 6: Condensed DiGraph

1.3 Simulation code

With the blocks of the model as given by the condensation of the model DiGraph, as previously discussed, we can generate *simulation code* for each block. That is *either*

- a symbolic function (definition) that takes exogenous input and returns the endogenous value (if the block is a definition) as described below, *or*

- a symbolic *objective function* and *Jacobian matrix* that takes exogenous input and (initial) values for the endogenous variables, which are to be sent to a Newton-Raphson algorithm.

A block is a said to be a definition *if and only if* it 1) consists of one equation and 2) the only thing on the left hand side of the equation is the endogenous variable of that equation (and that endogenous variable does *not* show up on the right hand side of that equation too).

1.4 Solution

In order to solve the model numerically, we loop (nested) over time periods (chronologically) and blocks (according to the order dictated by the condensation of the model DiGraph).

If the block to be solved is a definition, as discussed above, the solution is given by the function defining it (given exogenous and predetermined variables).

If the block to be solved is not a definition, it is sent to a Newton-Raphson algorithm, which in turn returns the solution (given exogenous and predetermined variables, and initial values for the endogenous variables). The $k + 1$st iteration of the Newton-Raphson algorithm is given by

$$\mathbf{x}_t^{(k+1)} = \mathbf{x}_t^{(k)} - \mathbf{J}_{\mathbf{F}}^{-1}(\mathbf{x}_t^{(k)}, \mathbf{z}_t)\mathbf{F}(\mathbf{x}_t^{(k)}, \mathbf{z}_t),$$

where $\mathbf{J}_{\mathbf{F}}(\mathbf{x}_t, \mathbf{z}_t)$ is the Jacobian matrix of $\mathbf{F}(\mathbf{x}_t, \mathbf{z}_t)$ evaluated at $\mathbf{x}_t$ and $\mathbf{z}_t$. The algorithm stops and returns $\mathbf{x}_t^{(k+1)}$ if $\max\left(\left|\mathbf{x}_t^{(k+1)} - \mathbf{x}_t^{(k)}\right|\right) \leq \varepsilon$, where

ε is a tolerance level. If $\mathbf{F}(\mathbf{x}_t^{(0)}, \mathbf{z}_t) = \mathbf{0}$ then the initial values solve the model, and as such $\mathbf{x}_t^{(0)}$ is returned as the solution.

2 Examples of use

The `ModelSolver` object is instantiated by

```
model = ModelSolver(equations, endogenous)
```

where `equations` and `endogenous` are lists containing equations and endogenous variables as strings, e.g.,

```
equations = [
'x1 = a1',
'x2 = a2',
'0.2*x1+0.7*x2 = 0.1*ca+0.8*cb+0.3*i1',
'0.8*x1+0.3*x2 = 0.9*ca+0.2*cb+0.1*i2',
'k1 = k1(-1)+i1',
'k2 = k2(-1)+i2'
]

endogenous = ['x1', 'x2', 'ca', 'cb', 'k1', 'k2']
```

Note that the `(-#)`-syntax denotes lags of a variable; `ModelSolver` does not support leads. Mathematical functions supported are `max()`, `min()`, `log()` and `exp()`. Note also that the equations are written in a form that is neither normalized, nor associates equations with endogenous variables.

It is possible to switch endogenous and exogenous variables using

```
model.switch_endo_vars(old_endo_vars, new_endo_vars)
```

where `old_endo_vars` and `new_endo_vars` are lists. Upon invoking `switch_endo_vars`, a tuple of endogenous variables stored inside the model object is updated, and `ModelSolver` immediately performs block analysis on the updated model, such that it can be solved for the updated list of endogenous variables.

2.1 Solving the model

Let `input_df` be a `Pandas DataFrame` containing exogenous variables and initial values for the endogenous variables. Then

```
solution_df = model.solve(input_df)
```

solves the model and stores the results in `solution_df`, which is a `Pandas DataFrame` identical to `input_df`, *but* where the endogenous variables are solutions to the model.

2.2 Analyzing the model

`ModelSolver` contains a number of methods that lets the user analyze the model.

`describe()` describes the model in terms of number of equations and blocks, along with information about the blocks, e.g.,

```
Model consists of 6 equations in 5 blocks
4 of the blocks consist of simple definitions

4 blocks have 1 equations
1 blocks have 2 equations
```

`find_endo_var(<endogenous variable>)` returns the block in which the endogenous variable is solved for.

`show_block(<block>)` returns information about the block, i.e., endogenous variables, exogenous and predetermined variables and equations.

`show_blocks()` returns information about all the blocks of the model.

`trace_to_exog_vars(<block>)` traces the model DiGraph back to the nodes of origin and reports what exogenous variables determine those nodes.

`trace_to_exog_vals(<block>, <period index>)` traces the model DiGraph back to the nodes of origin, finds what exogenous variables determine those nodes, and reports values for those variables for the chosen period index (0,1,...).

`draw_blockwise_graph()` draws a network graph as shown in Figure 7. The method takes as arguments a variable name, and the maximum number of ancestor and descendant nodes, and maximum number of nodes in total.

References

[1] Gilli, Manfred. "Causal Ordering and Beyond." International Economic Review, vol. 33, no. 4, 1992, pp. 957–71. JSTOR, https://doi.org/10.2307/2527152. Accessed 22 Mar. 2023.
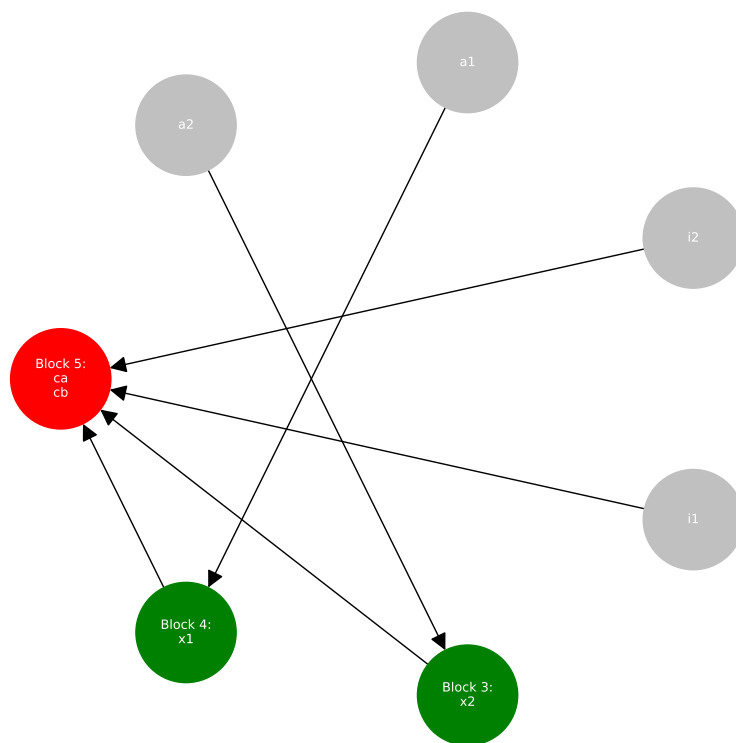
Figure 7: Example of network plot